## SEQUENCE LISTING

- <110> FURUKAWA, KEISUKE KAJIYAMA, NAOKI
- <120> MODIFIED SARCOSINE OXIDASES, GENES AND RECOMBINANT DNAS THEREOF, AND METHODS FOR PREPARING THE SAME
- <130> 252202US0
- <140> 10/829,427
- <141> 2004-04-22
- <150> JP 2003-121533
- <151> 2003-04-25
- <150> JP 2003-396807
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- <170> PatentIn version 3.3
- <210> 1
- <211> 387
- <212> PRT
- <213> Artificial Sequence
- <220>
- <223> Synthetic Peptide
- <400> 1
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- Thr Arg Ile Ile Arg His Ala Tyr Gly Glu Gly Arg Lys Tyr Val Pro 50 55 60

Phe Ala Leu Arg Ala Gln Glu Leu Trp Tyr Glu Leu Glu Asn Glu Thr His Asn Lys Ile Phe Thr Lys Thr Gly Val Leu Val Phe Gly Pro Lys Gly Glu Ser Asp Phe Val Ala Glu Thr Met Glu Ala Ala Glu His Ser Leu Thr Val Asp Leu Leu Glu Gly Asp Glu Ile Asn Thr Arg Trp Pro Gly Ile Thr Val Pro Glu Asn Tyr Asn Ala Ile Phe Glu Pro Asn Ser Gly Val Leu Phe Ser Glu Asn Cys Ile Arg Ser Tyr Arg Glu Leu Ala Val Ala Lys Gly Ala Lys Ile Leu Thr Tyr Thr Arg Val Glu Asp Phe Glu Val Ser Gln Asp Gln Val Lys Ile Gln Thr Ala Asn Gly Ser Tyr Thr Ala Asp Lys Leu Ile Val Ser Met Gly Ala Trp Asn Ser Lys Leu Leu Ser Lys Leu Asn Leu Asp Ile Pro Leu Gln Pro Tyr Arg Gln Val Val Gly Phe Phe Asp Ser Asn Glu Ala Lys Tyr Ser Asn Asp Val Gly Tyr Pro Ala Phe Met Val Glu Val Pro Lys Gly Ile Tyr Tyr Gly Phe Pro Ser Phe Gly Gly Cys Gly Leu Lys Ile Gly Tyr His Thr Tyr 

Gly Gln Gln Ile Asp Pro Asp Thr Ile Asn Arg Glu Phe Gly Ala Tyr 275 280 285 Gln Glu Asp Glu Ser Asn Leu Arg Asp Phe Leu Glu Lys Tyr Met Pro 300 295 Glu Ala Asn Gly Glu Leu Lys Arg Gly Ala Val Cys Met Tyr Thr Lys 315 320 310 305 Thr Pro Asp His His Phe Val Ile Asp Thr His Pro Glu His Ser Asn 330 325 Val Phe Val Ala Ala Gly Phe Ser Gly His Gly Phe Lys Phe Ser Ser 350 340 345 Val Val Gly Glu Val Leu Ser Gln Leu Ala Thr Thr Gly Lys Thr Glu 365 360 355 His Asp Ile Ser Ile Phe Ser Ile Asn Arg Pro Ala Leu Lys Gln Lys 380 375 370 Thr Thr Ile 385 <210> 2 1164 <211> <212> DNA Artificial Sequence <213> <220> <223> Synthetic DNA <400> atgagtacac attttgatgt gattgttgtt ggagcaggat caatgggaat ggctgcaggg tactatttag caaaacaagg agtcaaaaca ttattggtgg atgcattcga tccgccgcat acagaaggaa gccatcacgg tgatactcgc attatccgcc atgcttacgg tgaaggaaga

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60

120

180

240

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